

SEQUENCE LISTING

<110> Takeda Pharmaceutical Company Limited

<120> PREVENTIVES/REMEDIES FOR CANCER

<130> P04-222PCT

<150> JP 2002-240830

<151> 2002-08-21

<150> JP 2002-363108

<151> 2002-12-13

<150> PCT/JP2003/010532

<151> 2003-08-20

<160> 44

<210> 1

<211> 774

<212> PRT

<213> Homo sapiens

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25

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Arg Val Asp Cys Gly Gln Ala Pro Leu Asp Pro Val Tyr Leu His Val

35

40

45

Thr Ala Ala Arg Pro Ala Gln Pro Thr Leu Trp Thr Ala Lys Leu Asp

50

55

60

Arg Phe Lys Gly Ser Arg His His Thr Thr Leu Ile Thr Cys His Arg

65

70

75

80

Ala Gly Leu Thr Glu Pro Asp Ser Ser Ser Pro Leu Glu Leu Ser Glu

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Phe Leu Trp Val Asp Phe Val Val Glu Asn Ser Thr Gly Gly Gly Val		
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Ala Val Thr Arg Pro Val Thr Trp Gln Leu Glu Tyr Pro Gly Gln Ala		
115	120	125
Pro Glu Ala Glu Lys Asp Lys Met Val Trp Glu Ile Leu Val Ser Glu		
130	135	140
Arg Asp Ile Arg Ala Leu Ile Pro Leu Ala Lys Ala Glu Glu Leu Val		
145	150	155
Asn Thr Ala Pro Leu Thr Gly Val Pro Gln His Val Pro Val Arg Leu		
165	170	175
Val Thr Val Asp Gly Gly Gly Ala Leu Val Glu Val Thr Glu His Val		
180	185	190
Gly Cys Glu Ser Ala Asn Thr Gln Val Leu Gln Val Ser Glu Ala Cys		
195	200	205
Asp Ala Val Phe Val Ala Gly Lys Glu Ser Arg Gly Ala Arg Gly Val		
210	215	220
Arg Val Asp Phe Trp Trp Arg Arg Leu Arg Ala Ser Leu Arg Leu Thr		
225	230	235
Val Trp Ala Pro Leu Leu Pro Leu Arg Ile Glu Leu Thr Asp Thr Thr		
245	250	255
Leu Glu Gln Val Arg Gly Trp Arg Val Pro Gly Pro Ala Glu Gly Pro		
260	265	270
Ala Glu Pro Ala Ala Glu Ala Ser Asp Glu Ala Glu Arg Arg Ala Arg		
275	280	285
Gly Cys His Leu Gln Tyr Gln Arg Ala Gly Val Arg Phe Leu Ala Pro		
290	295	300
Phe Ala Ala His Pro Leu Asp Gly Gly Arg Arg Leu Thr His Leu Leu		
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Gly Pro Asp Trp Leu Leu Asp Val Ser His Leu Val Ala Pro His Ala		
325	330	335
Arg Val Leu Asp Ser Arg Val Ala Ser Leu Glu Gly Gly Arg Val Val		
340	345	350
Val Gly Arg Glu Pro Gly Val Thr Ser Ile Glu Val Arg Ser Pro Leu		
355	360	365
Ser Asp Ser Ile Leu Gly Glu Gln Ala Leu Ala Val Thr Asp Asp Lys		

370	375	380	
Val Ser Val Leu Glu Leu Arg Val Gln Pro Val Met Gly Ile Ser Leu			
385	390	395	400
Thr Leu Ser Arg Gly Thr Ala His Pro Gly Glu Val Thr Ala Thr Cys			
	405	410	415
Trp Ala Gln Ser Ala Leu Pro Ala Pro Lys Gln Glu Val Ala Leu Ser			
	420	425	430
Leu Trp Leu Ser Phe Ser Asp His Thr Val Ala Pro Ala Glu Leu Tyr			
	435	440	445
Asp Arg Arg Asp Leu Gly Leu Ser Val Ser Ala Glu Glu Pro Gly Ala			
	450	455	460
Ile Leu Pro Ala Glu Glu Gln Gly Ala Gln Leu Gly Val Val Val Ser			
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Gly Ala Gly Ala Glu Gly Leu Pro Leu His Val Ala Leu His Pro Pro			
	485	490	495
Glu Pro Cys Arg Arg Gly Arg His Arg Val Pro Leu Ala Ser Gly Thr			
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Ala Trp Leu Gly Leu Pro Pro Ala Ser Thr Pro Ala Pro Ala Leu Pro			
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Lys Arg Gln Val Ala Gly Ser Val Gly Gly Asn Thr Gly Val Arg Gly			
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Lys Phe Glu Arg Ala Glu Glu Glu Ala Arg Lys Glu Glu Thr Glu Ala			
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Arg Glu Glu Glu Glu Glu Glu Glu Glu Met Val Pro Ala Pro Gln			
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His Val Thr Glu Leu Glu Leu Gly Met Tyr Ala Leu Leu Gly Val Phe			
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Cys Val Ala Ile Phe Ile Phe Leu Val Asn Gly Val Val Phe Val Leu			
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Arg Tyr Gln Arg Lys Glu Pro Pro Asp Ser Ala Thr Asp Pro Thr Ser			
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Pro Gln Pro His Asn Trp Val Trp Leu Gly Thr Asp Gln Glu Glu Leu			
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Ser Arg Gln Leu Asp Arg Gln Ser Pro Gly Pro Pro Lys Gly Glu Gly			

660	665	670
Ser Cys Pro Cys Glu Ser Gly Gly Gly Glu Ala Pro Thr Leu Ala		
675	680	685
Pro Gly Pro Pro Gly Gly Thr Thr Ser Ser Ser Ser Thr Leu Ala Arg		
690	695	700
Lys Glu Ala Gly Gly Arg Arg Lys Arg Val Glu Phe Val Thr Phe Val		
705	710	715
Pro Ala Pro Pro Ala Gln Ser Pro Glu Glu Pro Val Gly Ala Pro Ala		
725	730	735
Val Gln Ser Ile Leu Val Ala Gly Glu Glu Asp Ile Arg Trp Val Cys		
740	745	750
Glu Asp Met Gly Leu Lys Asp Pro Glu Glu Leu Arg Asn Tyr Met Glu		
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<210> 2

<211> 2322

<212> DNA

<213> Homo sapiens

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<210> 3

<211> 2755

<212> DNA

<213> Homo sapiens

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<210> 4

<211> 909

<212> PRT

<213> Homo sapiens

<400> 4

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Arg Val Asp Cys Gly Gln Ala Pro Leu Asp Pro Val Tyr Leu Pro Ala
      35              40              45
Ala Leu Glu Leu Leu Asp Ala Pro Glu His Phe Arg Val Gln Gln Val
      50              55              60
Gly His Tyr Pro Pro Ala Asn Ser Ser Leu Ser Ser Arg Ser Glu Thr
      65              70              75              80
Phe Leu Leu Leu Gln Pro Trp Pro Arg Ala Gln Pro Leu Leu Arg Ala
      85              90              95
Ser Tyr Pro Pro Phe Ala Thr Gln Gln Val Val Pro Pro Arg Val Thr
      100             105             110
Glu Pro His Gln Arg Pro Val Pro Trp Asp Val Arg Ala Val Ser Val
      115             120             125
Glu Ala Ala Val Thr Pro Ala Glu Pro Tyr Ala Arg Val Leu Phe His
      130             135             140
Leu Lys Gly Gln Asp Trp Pro Pro Gly Ser Gly Ser Leu Pro Cys Ala
      145             150             155             160
Arg Leu His Ala Thr His Pro Ala Gly Thr Ala His Gln Ala Cys Arg
      165             170             175
Phe Gln Pro Ser Leu Gly Ala Cys Val Val Glu Leu Glu Leu Pro Ser

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Leu Glu Pro Ala Ala Glu Gly Pro Gly Gly Cys Gly Ser Gly Glu Glu		
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Asn Asp Pro Gly Glu Gln Ala Leu Pro Val Gly Gly Val Glu Leu Arg		
225	230	235
Pro Ala Asp Pro Pro Gln Tyr Gln Glu Val Pro Leu Asp Glu Ala Val		
245	250	255
Thr Leu Arg Val Pro Asp Met Pro Val Arg Pro Gly Gln Leu Phe Ser		
260	265	270
Ala Thr Leu Leu Leu Arg His Asn Phe Thr Ala Ser Leu Leu Thr Leu		
275	280	285
Arg Ile Lys Val Lys Lys Gly Leu His Val Thr Ala Ala Arg Pro Ala		
290	295	300
Gln Pro Thr Leu Trp Thr Ala Lys Leu Asp Arg Phe Lys Gly Ser Arg		
305	310	315
His His Thr Thr Leu Ile Thr Cys His Arg Ala Gly Leu Thr Glu Pro		
325	330	335
Asp Ser Ser Pro Leu Glu Leu Ser Glu Phe Leu Trp Val Asp Phe Val		
340	345	350
Val Glu Asn Ser Thr Gly Gly Gly Val Ala Val Thr Arg Pro Val Thr		
355	360	365
Trp Gln Leu Glu Tyr Pro Gly Gln Ala Pro Glu Ala Glu Lys Asp Lys		
370	375	380
Met Val Trp Glu Ile Leu Val Ser Glu Arg Asp Ile Arg Ala Leu Ile		
385	390	395
Pro Leu Ala Lys Ala Glu Glu Leu Val Asn Thr Ala Pro Leu Thr Gly		
405	410	415
Val Pro Gln His Val Pro Val Arg Leu Val Thr Val Asp Gly Gly Gly		
420	425	430
Ala Leu Val Glu Val Thr Glu His Val Gly Cys Glu Ser Ala Asn Thr		
435	440	445
Gln Val Leu Gln Val Ser Glu Ala Cys Asp Ala Val Phe Val Ala Gly		
450	455	460
Lys Glu Ser Arg Gly Ala Arg Gly Val Arg Val Asp Phe Trp Trp Arg		

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Leu Arg Ile Glu Leu Thr Asp Thr Thr Leu Glu Gln Val Arg Gly Trp			
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Arg Val Pro Gly Pro Ala Glu Gly Pro Ala Glu Pro Ala Ala Glu Ala			
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Ser Asp Glu Ala Glu Arg Arg Ala Arg Gly Cys His Leu Gln Tyr Gln			
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Arg Ala Gly Val Arg Phe Leu Ala Pro Phe Ala Ala His Pro Leu Asp			
545	550	555	560
Gly Gly Arg Arg Leu Thr His Leu Leu Gly Pro Asp Trp Leu Leu Asp			
565	570	575	
Val Ser His Leu Val Ala Pro His Ala Arg Val Leu Asp Ser Arg Val			
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Thr Ser Ile Glu Val Arg Ser Pro Leu Ser Asp Ser Ile Leu Gly Glu			
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Gln Ala Leu Ala Val Thr Asp Asp Lys Val Ser Val Leu Glu Leu Arg			
625	630	635	640
Val Gln Pro Val Met Gly Ile Ser Leu Thr Leu Ser Arg Gly Thr Ala			
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His Pro Gly Glu Val Thr Ala Thr Cys Trp Ala Gln Ser Ala Leu Pro			
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Ala Pro Lys Gln Glu Val Ala Leu Ser Leu Trp Leu Ser Phe Ser Asp			
675	680	685	
His Thr Val Ala Pro Ala Glu Leu Tyr Asp Arg Arg Asp Leu Gly Leu			
690	695	700	
Ser Val Ser Ala Glu Glu Pro Gly Ala Ile Leu Pro Ala Glu Glu Gln			
705	710	715	720
Gly Ala Gln Leu Gly Val Val His Val Thr Glu Leu Glu Leu Gly Met			
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Tyr Ala Leu Leu Gly Val Phe Cys Val Ala Ile Phe Ile Phe Leu Val			
740	745	750	
Asn Gly Val Val Phe Val Leu Arg Tyr Gln Arg Lys Glu Pro Pro Asp			

755	760	765
Ser Ala Thr Asp Pro Thr Ser Pro Gln Pro His Asn Trp Val Trp Leu		
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Gly Thr Asp Gln Glu Glu Leu Ser Arg Gln Leu Asp Arg Gln Ser Pro		
785	790	795
Gly Pro Pro Lys Gly Glu Gly Ser Cys Pro Cys Glu Ser Gly Gly Gly		
805	810	815
Gly Glu Ala Pro Thr Leu Ala Pro Gly Pro Pro Gly Gly Thr Thr Ser		
820	825	830
Ser Ser Ser Thr Leu Ala Arg Lys Glu Ala Gly Gly Arg Arg Lys Arg		
835	840	845
Val Glu Phe Val Thr Phe Ala Pro Ala Pro Pro Ala Gln Ser Pro Glu		
850	855	860
Glu Pro Val Gly Ala Pro Ala Val Gln Ser Ile Leu Val Ala Gly Glu		
865	870	875
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Glu Leu Arg Asn Tyr Met Glu Arg Ile Arg Gly Ser Ser		
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<210> 5

<211> 2727

<212> DNA

<213> Homo sapiens

<400> 5

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<210> 6

<211> 2778

<212> DNA

<213> Homo sapiens

<400> 6

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<210> 7

<211> 594

<212> PRT

<213> Homo sapiens

<400> 7

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Arg Val Asp Cys Gly Gln Ala Pro Leu Asp Pro Gly Leu His Val Thr
      35              40              45
Ala Ala Arg Pro Ala Gln Pro Thr Leu Trp Thr Ala Lys Leu Asp Arg
      50              55              60
Phe Lys Gly Ser Arg His His Thr Thr Leu Ile Thr Cys His Arg Ala
      65              70              75              80
Gly Leu Thr Glu Pro Asp Ser Ser Ser Pro Leu Glu Leu Ser Glu Phe

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	85		90		95
Leu Trp Val Asp Phe Val Val Glu Asn Ser Thr Gly Gly Gly Val Ala					
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Val Thr Arg Pro Val Thr Trp Gln Leu Glu Tyr Pro Gly Gln Ala Pro					
115		120		125	
Glu Ala Glu Lys Asp Lys Met Val Trp Glu Ile Leu Val Ser Glu Arg					
130		135		140	
Asp Ile Arg Ala Leu Ile Pro Leu Ala Lys Ala Glu Glu Leu Val Asn					
145		150		155	160
Thr Ala Pro Leu Thr Gly Val Pro Gln His Val Pro Val Arg Leu Val					
	165		170		175
Thr Val Asp Gly Gly Gly Ala Leu Val Glu Val Thr Glu His Val Gly					
	180		185		190
Cys Glu Ser Ala Asn Thr Gln Val Leu Gln Val Ser Glu Ala Cys Asp					
	195		200		205
Ala Val Phe Val Ala Gly Lys Glu Ser Arg Gly Ala Arg Gly Val Arg					
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Val Asp Phe Trp Trp Arg Arg Leu Arg Ala Ser Leu Arg Leu Thr Val					
225		230		235	240
Trp Ala Pro Leu Leu Pro Leu Arg Ile Glu Leu Thr Asp Thr Thr Leu					
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Glu Gln Val Arg Gly Trp Arg Val Pro Gly Pro Ala Glu Gly Pro Ala					
	260		265		270
Glu Pro Ala Ala Glu Ala Ser Asp Glu Ala Glu Arg Arg Ala Arg Gly					
	275		280		285
Cys His Leu Gln Tyr Gln Arg Ala Gly Val Arg Phe Leu Ala Pro Phe					
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Ala Ala His Pro Leu Asp Gly Gly Arg Arg Leu Thr His Leu Leu Gly					
305		310		315	320
Pro Asp Trp Leu Leu Asp Val Ser His Leu Val Ala Pro His Ala Arg					
	325		330		335
Val Leu Asp Ser Arg Val Ala Ser Leu Glu Gly Gly Arg Val Val Val					
	340		345		350
Gly Arg Glu Pro Gly Val Thr Ser Ile Glu Val Arg Ser Pro Leu Ser					
	355		360		365
Asp Ser Ile Leu Gly Glu Gln Ala Leu Ala Val Thr Asp Asp Lys Val					

370	375	380	
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385	390	395	400
Leu Ser Arg Gly Thr Ala His Pro Gly Glu Val Thr Ala Thr Cys Trp			
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Ala Gln Ser Ala Leu Pro Ala Pro Lys Gln Glu Val Ala Leu Ser Leu			
	420	425	430
Trp Leu Ser Phe Ser Asp His Thr Val Ala Pro Ala Glu Leu Tyr Asp			
	435	440	445
Arg Arg Asp Leu Gly Leu Ser Val Ser Ala Glu Glu Pro Gly Ala Ile			
	450	455	460
Leu Pro Ala Glu Glu Gln Gly Ala Gln Leu Gly Val Val Val Ser Gly			
465	470	475	480
Ala Gly Ala Glu Gly Leu Pro Leu His Val Ala Leu His Pro Pro Glu			
	485	490	495
Pro Cys Arg Arg Gly Arg His Arg Val Pro Leu Ala Ser Gly Thr Ala			
	500	505	510
Trp Leu Gly Leu Pro Pro Ala Ser Thr Pro Ala Pro Ala Leu Pro Ser			
	515	520	525
Ser Pro Ala Trp Ser Pro Pro Ala Thr Glu Ala Thr Met Gly Gly Lys			
	530	535	540
Arg Gln Val Ala Gly Ser Val Gly Gly Asn Thr Gly Val Arg Gly Lys			
545	550	555	560
Phe Glu Arg Ala Glu Glu Glu Ala Arg Lys Glu Glu Thr Glu Ala Arg			
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Asp Gly Gly Gly Gly Arg Gly Gly Gly Asp Gly Pro Cys Pro Ser Ala			
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Cys His			

<210> 8

<211> 1782

<212> DNA

<213> Homo sapiens

<400> 8

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<210> 9

<211> 2735

<212> DNA

<213> Homo sapiens

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<210> 10

<211> 639

<212> PRT

<213> Homo sapiens

<400> 10

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      20              25              30
Val Pro Gln His Val Pro Val Arg Leu Val Thr Val Asp Gly Gly Gly
      35              40              45
Ala Leu Val Glu Val Thr Glu His Val Gly Cys Glu Ser Ala Asn Thr
      50              55              60
Gln Val Leu Gln Val Ser Glu Ala Cys Asp Ala Val Phe Val Ala Gly
      65              70              75              80
Lys Glu Ser Arg Gly Ala Arg Gly Val Arg Val Asp Phe Trp Trp Arg
      85              90              95
Arg Leu Arg Ala Ser Leu Arg Leu Thr Val Trp Ala Pro Leu Leu Pro
      100             105             110
Leu Arg Ile Glu Leu Thr Asp Thr Thr Leu Glu Gln Val Arg Gly Trp
      115             120             125
Arg Val Pro Gly Pro Ala Glu Gly Pro Ala Glu Pro Ala Ala Glu Ala
      130             135             140
Ser Asp Glu Ala Glu Arg Arg Ala Arg Gly Cys His Leu Gln Tyr Gln

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165	170	175	
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180	185	190	
Val Ser His Leu Val Ala Pro His Ala Arg Val Leu Asp Ser Arg Val			
195	200	205	
Ala Ser Leu Glu Gly Gly Arg Val Val Val Gly Arg Glu Pro Gly Val			
210	215	220	
Thr Ser Ile Glu Val Arg Ser Pro Leu Ser Asp Ser Ile Leu Gly Glu			
225	230	235	240
Gln Ala Leu Ala Val Thr Asp Asp Lys Val Ser Val Leu Glu Leu Arg			
245	250	255	
Val Gln Pro Val Met Gly Ile Ser Leu Thr Leu Ser Arg Gly Thr Ala			
260	265	270	
His Pro Gly Glu Val Thr Ala Thr Cys Trp Ala Gln Ser Ala Leu Pro			
275	280	285	
Ala Pro Lys Gln Glu Val Ala Leu Ser Leu Trp Leu Ser Phe Ser Asp			
290	295	300	
His Thr Val Ala Pro Ala Glu Leu Tyr Asp Arg Arg Asp Leu Gly Leu			
305	310	315	320
Ser Val Ser Ala Glu Glu Pro Gly Ala Ile Leu Pro Ala Glu Glu Gln			
325	330	335	
Gly Ala Gln Leu Gly Val Val Val Ser Gly Ala Gly Ala Glu Gly Leu			
340	345	350	
Pro Leu His Val Ala Leu His Pro Pro Glu Pro Cys Arg Arg Gly Arg			
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His Arg Val Pro Leu Ala Ser Gly Thr Ala Trp Leu Gly Leu Pro Pro			
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Ala Ser Thr Pro Ala Pro Ala Leu Pro Ser Ser Pro Ala Trp Ser Pro			
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Pro Ala Thr Glu Ala Thr Met Gly Gly Lys Arg Gln Val Ala Gly Ser			
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Val Gly Gly Asn Thr Gly Val Arg Gly Lys Phe Glu Arg Ala Glu Glu			
420	425	430	
Glu Ala Arg Lys Glu Glu Thr Lys Ala Arg Glu Glu Glu Glu Glu			

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Gly Met Tyr Ala Leu Leu Gly Val Phe Cys Val Ala Ile Phe Ile Phe		
465	470	475
Leu Val Asn Gly Val Val Phe Val Leu Arg Tyr Gln Arg Lys Glu Pro		
485	490	495
Pro Asp Ser Ala Thr Asp Pro Thr Ser Pro Gln Pro His Asn Trp Val		
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Trp Leu Gly Thr Asp Gln Glu Glu Leu Ser Arg Gln Leu Asp Arg Gln		
515	520	525
Ser Pro Gly Pro Pro Lys Gly Glu Gly Ser Cys Pro Cys Glu Ser Gly		
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Gly Gly Gly Glu Ala Pro Thr Leu Ala Pro Gly Pro Pro Gly Gly Thr		
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Thr Ser Ser Ser Ser Thr Leu Ala Arg Lys Glu Ala Gly Gly Arg Arg		
565	570	575
Lys Arg Val Glu Phe Val Thr Phe Ala Pro Ala Pro Pro Ala Gln Ser		
580	585	590
Pro Glu Glu Pro Val Gly Ala Pro Ala Val Gln Ser Ile Leu Val Ala		
595	600	605
Gly Glu Glu Asp Ile Arg Trp Val Cys Glu Asp Met Gly Leu Lys Asp		
610	615	620
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625	630	635

<210> 11

<211> 1917

<212> DNA

<213> Homo sapiens

<400> 11

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<211> 2235

<212> DNA

<213> Homo sapiens

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2235

<210> 13

<211> 20

<212> DNA

<213> Artificial Sequence

<220>

<223> Designed oligonucleotide

<400> 13

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<210> 14

<211> 20

<212> DNA

<213> Artificial Sequence

<220>

<223> Designed oligonucleotide

<400> 14

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<210> 15

<211> 1024

<212> PRT

<213> Homo sapiens

<400> 15

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5

10

15

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20

25

30

Arg Val Asp Cys Gly Gln Ala Pro Leu Asp Pro Val Tyr Leu Pro Ala

35

40

45

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 85 90 95
 Ser Tyr Pro Pro Phe Ala Thr Gln Gln Val Val Pro Pro Arg Val Thr
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 Glu Pro His Gln Arg Pro Val Pro Trp Asp Val Arg Ala Val Ser Val
 115 120 125
 Glu Ala Ala Val Thr Pro Ala Glu Pro Tyr Ala Arg Val Leu Phe His
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 Leu Lys Gly Gln Asp Trp Pro Pro Gly Ser Gly Ser Leu Pro Cys Ala
 145 150 155 160
 Arg Leu His Ala Thr His Pro Ala Gly Thr Ala His Gln Ala Cys Arg
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 Phe Gln Pro Ser Leu Gly Ala Cys Val Val Glu Leu Glu Leu Pro Ser
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 His Trp Phe Ser Gln Ala Ser Thr Thr Arg Ala Glu Leu Ala Tyr Thr
 195 200 205
 Leu Glu Pro Ala Ala Glu Gly Pro Gly Gly Cys Gly Ser Gly Glu Glu
 210 215 220
 Asn Asp Pro Gly Glu Gln Ala Leu Pro Val Gly Gly Val Glu Leu Arg
 225 230 235 240
 Pro Ala Asp Pro Pro Gln Tyr Gln Glu Val Pro Leu Asp Glu Ala Val
 245 250 255
 Thr Leu Arg Val Pro Asp Met Pro Val Arg Pro Gly Gln Leu Phe Ser
 260 265 270
 Ala Thr Leu Leu Leu Arg His Asn Phe Thr Ala Ser Leu Leu Thr Leu
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 Arg Ile Lys Val Lys Lys Gly Leu His Val Thr Ala Ala Arg Pro Ala
 290 295 300
 Gln Pro Thr Leu Trp Thr Ala Lys Leu Asp Arg Phe Lys Gly Ser Arg
 305 310 315 320
 His His Thr Thr Leu Ile Thr Cys His Arg Ala Gly Leu Thr Glu Pro
 325 330 335

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Val Val Glu Asn Ser Thr Gly Gly Gly Val Ala Val Thr Arg Pro Val			
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Thr Trp Gln Leu Glu Tyr Pro Gly Gln Ala Pro Glu Ala Glu Lys Asp			
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Lys Met Val Trp Glu Ile Leu Val Ser Glu Arg Asp Ile Arg Ala Leu			
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Ile Pro Leu Ala Lys Ala Glu Glu Leu Val Asn Thr Ala Pro Leu Thr			
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Gly Val Pro Gln His Val Pro Val Arg Leu Val Thr Val Asp Gly Gly			
420	425	430	
Gly Ala Leu Val Glu Val Thr Glu His Val Gly Cys Glu Ser Ala Asn			
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Thr Gln Val Leu Gln Val Ser Glu Ala Cys Asp Ala Val Phe Val Ala			
450	455	460	
Gly Lys Glu Ser Arg Gly Ala Arg Gly Val Arg Val Asp Phe Trp Trp			
465	470	475	480
Arg Arg Leu Arg Ala Ser Leu Arg Leu Thr Val Trp Ala Pro Leu Leu			
485	490	495	
Pro Leu Arg Ile Glu Leu Thr Asp Thr Thr Leu Glu Gln Val Arg Gly			
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Trp Arg Val Pro Gly Pro Ala Glu Gly Pro Ala Glu Pro Ala Ala Glu			
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Gln Arg Ala Gly Val Arg Phe Leu Ala Pro Phe Ala Ala His Pro Leu			
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Asp Gly Gly Arg Arg Leu Thr His Leu Leu Gly Pro Asp Trp Leu Leu			
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Asp Val Ser His Leu Val Ala Pro His Ala Arg Val Leu Asp Ser Arg			
580	585	590	
Val Ala Ser Leu Glu Gly Gly Arg Val Val Val Gly Arg Glu Pro Gly			
595	600	605	
Val Thr Ser Ile Glu Val Arg Ser Pro Leu Ser Asp Ser Ile Leu Gly			
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Glu Gln Ala Leu Ala Val Thr Asp Asp Lys Val Ser Val Leu Glu Leu			
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Arg Val Gln Pro Val Met Gly Ile Ser Leu Thr Leu Ser Arg Gly Thr			
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Ala His Pro Gly Glu Val Thr Ala Thr Cys Trp Ala Gln Ser Ala Leu			
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Pro Ala Pro Lys Gln Glu Val Ala Leu Ser Leu Trp Leu Ser Phe Ser			
	675	680	685
Asp His Thr Val Ala Pro Ala Glu Leu Tyr Asp Arg Arg Asp Leu Gly			
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Leu Ser Val Ser Ala Glu Glu Pro Gly Ala Ile Leu Pro Ala Glu Glu			
705	710	715	720
Gln Gly Ala Gln Leu Gly Val Val Val Ser Gly Ala Gly Ala Glu Gly			
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Leu Pro Leu His Val Ala Leu His Pro Pro Glu Pro Cys Arg Arg Gly			
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Arg His Arg Val Pro Leu Ala Ser Gly Thr Ala Trp Leu Gly Leu Pro			
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Pro Ala Ser Thr Pro Ala Pro Ala Leu Pro Ser Ser Pro Ala Trp Ser			
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Pro Pro Ala Thr Glu Ala Thr Met Gly Gly Lys Arg Gln Val Ala Gly			
785	790	795	800
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	805	810	815
Glu Glu Ala Arg Lys Glu Glu Thr Glu Ala Arg Glu Glu Glu Glu Glu			
	820	825	830
Glu Glu Glu Glu Met Val Pro Ala Pro Gln His Val Thr Glu Leu Glu			
	835	840	845
Leu Gly Met Tyr Ala Leu Leu Gly Val Phe Cys Val Ala Ile Phe Ile			
	850	855	860
Phe Leu Val Asn Gly Val Val Phe Val Leu Arg Tyr Gln Arg Lys Glu			
865	870	875	880
Pro Pro Asp Ser Ala Thr Asp Pro Thr Ser Pro Gln Pro His Asn Trp			
	885	890	895
Val Trp Leu Gly Thr Asp Gln Glu Glu Leu Ser Arg Gln Leu Asp Arg			
	900	905	910

Gln Ser Pro Gly Pro Pro Lys Gly Glu Gly Ser Cys Pro Cys Glu Ser
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 Gly Gly Gly Gly Glu Ala Pro Thr Leu Ala Pro Gly Pro Pro Gly Gly
 930 935 940
 Thr Thr Ser Ser Ser Ser Thr Leu Ala Arg Lys Glu Ala Gly Gly Arg
 945 950 955 960
 Arg Lys Arg Val Glu Phe Val Thr Phe Val Pro Ala Pro Pro Ala Gln
 965 970 975
 Ser Pro Glu Glu Pro Val Gly Ala Pro Ala Val Gln Ser Ile Leu Val
 980 985 990
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<210> 16

<211> 3072

<212> DNA

<213> Homo sapiens

<400> 16

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3072

<210> 17

<211> 1020

<212> PRT

<213> Homo sapiens

<400> 17

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      35              40              45
Leu Asp Ala Pro Glu His Phe Arg Val Gln Gln Val Gly His Tyr Pro
      50              55              60
Pro Ala Asn Ser Ser Leu Ser Ser Arg Ser Glu Thr Phe Leu Leu Leu
      65              70              75              80
Gln Pro Trp Pro Arg Ala Gln Pro Leu Leu Arg Ala Ser Tyr Pro Pro
      85              90              95
Phe Ala Thr Gln Gln Val Val Pro Pro Arg Val Thr Glu Pro His Gln
      100             105             110
Arg Pro Val Pro Trp Asp Val Arg Ala Val Ser Val Glu Ala Ala Val
      115             120             125
Thr Pro Ala Glu Pro Tyr Ala Arg Val Leu Phe His Leu Lys Gly Gln
      130             135             140
Asp Trp Pro Pro Gly Ser Gly Ser Leu Pro Cys Ala Arg Leu His Ala
      145             150             155             160
Thr His Pro Ala Gly Thr Ala His Gln Ala Cys Arg Phe Gln Pro Ser
      165             170             175
Leu Gly Ala Cys Val Val Glu Leu Glu Leu Pro Ser His Trp Phe Ser
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Gln Ala Ser Thr Thr Arg Ala Glu Leu Ala Tyr Thr Leu Glu Pro Ala
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Ala Glu Gly Pro Gly Gly Cys Gly Ser Gly Glu Glu Asn Asp Pro Gly
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 Arg Leu Thr His Leu Leu Gly Pro Asp Trp Leu Leu Asp Val Ser His
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 595 600 605
 Glu Val Arg Ser Pro Leu Ser Asp Ser Ile Leu Gly Glu Gln Ala Leu
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 Ala Val Thr Asp Asp Lys Val Ser Val Leu Glu Leu Arg Val Gln Pro
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 Val Met Gly Ile Ser Leu Thr Leu Ser Arg Gly Thr Ala His Pro Gly
 645 650 655
 Glu Val Thr Ala Thr Cys Trp Ala Gln Ser Ala Leu Pro Ala Pro Lys
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 Gln Glu Val Ala Leu Ser Leu Trp Leu Ser Phe Ser Asp His Thr Val
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 740 745 750
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 Glu Ala Thr Met Gly Gly Lys Arg Gln Val Ala Gly Ser Val Gly Gly
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Lys Glu Glu Thr Glu Ala Arg Glu Glu Glu Glu Glu Glu Glu Glu			
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Ala Leu Leu Gly Val Phe Cys Val Ala Ile Phe Ile Phe Leu Val Asn			
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Gly Val Val Phe Val Leu Arg Tyr Gln Arg Lys Glu Pro Pro Asp Ser			
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Ala Thr Asp Pro Thr Ser Pro Gln Pro His Asn Trp Val Trp Leu Gly			
885	890	895	
Thr Asp Gln Glu Glu Leu Ser Arg Gln Leu Asp Arg Gln Ser Pro Gly			
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Pro Pro Lys Gly Glu Gly Ser Cys Pro Cys Glu Ser Gly Gly Gly Gly			
915	920	925	
Glu Ala Pro Thr Leu Ala Pro Gly Pro Pro Gly Gly Thr Thr Ser Ser			
930	935	940	
Ser Ser Thr Leu Ala Arg Lys Glu Ala Gly Gly Arg Arg Lys Arg Val			
945	950	955	960
Glu Phe Val Thr Phe Val Pro Ala Pro Pro Ala Gln Ser Pro Glu Glu			
965	970	975	
Pro Val Gly Ala Pro Ala Val Gln Ser Ile Leu Val Ala Gly Glu Glu			
980	985	990	
Asp Ile Arg Trp Val Cys Glu Asp Met Gly Leu Lys Asp Pro Glu Glu			
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<210> 18

<211> 3060

<212> DNA

<213> Homo sapiens

<400> 18

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<211> 3505

<212> DNA

<213> Homo sapiens

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<210> 20

<211> 1023

<212> PRT

<213> Homo sapiens

<400> 20

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Arg Val Asp Cys Gly Gln Ala Pro Leu Asp Pro Val Tyr Leu Pro Ala
      35              40              45
Ala Leu Glu Leu Leu Asp Ala Pro Glu His Phe Arg Val Gln Gln Val
      50              55              60
Gly His Tyr Pro Pro Ala Asn Ser Ser Leu Ser Ser Arg Ser Glu Thr
      65              70              75              80
Phe Leu Leu Leu Gln Pro Trp Pro Arg Ala Gln Pro Leu Leu Arg Ala
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Ser Tyr Pro Pro Phe Ala Thr Gln Gln Val Val Pro Pro Arg Val Thr
      100             105             110
Glu Pro His Gln Arg Pro Val Pro Trp Asp Val Arg Ala Val Ser Val
      115             120             125
Glu Ala Ala Val Thr Pro Ala Glu Pro Tyr Ala Arg Val Leu Phe His
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Leu Lys Gly Gln Asp Trp Pro Pro Gly Ser Gly Ser Leu Pro Cys Ala
      145             150             155             160
Arg Leu His Ala Thr His Pro Ala Gly Thr Ala His Gln Ala Cys Arg

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His	Trp	Phe	Ser	Gln	Ala	Ser	Thr	Thr	Arg	Ala	Glu	Leu	Ala	Tyr	Thr
	195				200					205					
Leu	Glu	Pro	Ala	Ala	Glu	Gly	Pro	Gly	Gly	Cys	Gly	Ser	Gly	Glu	Glu
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Asn	Asp	Pro	Gly	Glu	Gln	Ala	Leu	Pro	Val	Gly	Gly	Val	Glu	Leu	Arg
225				230						235				240	
Pro	Ala	Asp	Pro	Pro	Gln	Tyr	Gln	Glu	Val	Pro	Leu	Asp	Glu	Ala	Val
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Arg	Ile	Lys	Val	Lys	Lys	Gly	Leu	His	Val	Thr	Ala	Ala	Arg	Pro	Ala
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Asp	Ser	Ser	Pro	Leu	Glu	Leu	Ser	Glu	Phe	Leu	Trp	Val	Asp	Phe	Val
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Val	Glu	Asn	Ser	Thr	Gly	Gly	Gly	Val	Ala	Val	Thr	Arg	Pro	Val	Thr
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Trp	Gln	Leu	Glu	Tyr	Pro	Gly	Gln	Ala	Pro	Glu	Ala	Glu	Lys	Asp	Lys
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Gln	Val	Leu	Gln	Val	Ser	Glu	Ala	Cys	Asp	Ala	Val	Phe	Val	Ala	Gly

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485	490	495	
Leu Arg Ile Glu Leu Thr Asp Thr Thr Leu Glu Gln Val Arg Gly Trp			
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Arg Val Pro Gly Pro Ala Glu Gly Pro Ala Glu Pro Ala Ala Glu Ala			
515	520	525	
Ser Asp Glu Ala Glu Arg Arg Ala Arg Gly Cys His Leu Gln Tyr Gln			
530	535	540	
Arg Ala Gly Val Arg Phe Leu Ala Pro Phe Ala Ala His Pro Leu Asp			
545	550	555	560
Gly Gly Arg Arg Leu Thr His Leu Leu Gly Pro Asp Trp Leu Leu Asp			
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His Pro Gly Glu Val Thr Ala Thr Cys Trp Ala Gln Ser Ala Leu Pro			
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Ala Pro Lys Gln Glu Val Ala Leu Ser Leu Trp Leu Ser Phe Ser Asp			
675	680	685	
His Thr Val Ala Pro Ala Glu Leu Tyr Asp Arg Arg Asp Leu Gly Leu			
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Ser Val Ser Ala Glu Glu Pro Gly Ala Ile Leu Pro Ala Glu Glu Gln			
705	710	715	720
Gly Ala Gln Leu Gly Val Val Val Ser Gly Ala Gly Ala Glu Gly Leu			
725	730	735	
Pro Leu His Val Ala Leu His Pro Pro Glu Pro Cys Arg Arg Gly Arg			

740	745	750
His Arg Val Pro Leu Ala Ser Gly Thr Ala Trp Leu Gly Leu Pro Pro		
755	760	765
Ala Ser Thr Pro Ala Pro Ala Leu Pro Ser Ser Pro Ala Trp Ser Pro		
770	775	780
Pro Ala Thr Glu Ala Thr Met Gly Gly Lys Arg Gln Val Ala Gly Ser		
785	790	795
Val Gly Gly Asn Thr Gly Val Arg Gly Lys Phe Glu Arg Ala Glu Glu		
805	810	815
Glu Ala Arg Lys Glu Glu Thr Lys Pro Arg Glu Glu Glu Glu Glu Glu		
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Glu Glu Glu Met Val Pro Ala Pro Gln His Val Thr Glu Leu Glu Leu		
835	840	845
Gly Met Tyr Ala Leu Leu Gly Val Phe Cys Val Ala Ile Phe Ile Phe		
850	855	860
Leu Val Asn Gly Val Val Phe Val Leu Arg Tyr Gln Arg Lys Glu Pro		
865	870	875
Pro Asp Ser Ala Thr Asp Pro Thr Ser Pro Gln Pro His Asn Trp Val		
885	890	895
Trp Leu Gly Thr Asp Gln Glu Glu Leu Ser Arg Gln Leu Asp Arg Gln		
900	905	910
Ser Pro Gly Pro Pro Lys Gly Glu Gly Ser Cys Pro Cys Glu Ser Gly		
915	920	925
Gly Gly Gly Glu Ala Pro Thr Leu Ala Pro Gly Pro Pro Gly Gly Thr		
930	935	940
Thr Ser Ser Ser Ser Thr Leu Ala Arg Lys Glu Ala Gly Gly Arg Arg		
945	950	955
Lys Arg Val Glu Phe Val Thr Phe Ala Pro Ala Pro Pro Ala Gln Ser		
965	970	975
Pro Glu Glu Pro Val Gly Ala Pro Ala Val Gln Ser Ile Leu Val Ala		
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<210> 21

<211> 3069

<212> DNA

<213> Homo sapiens

<400> 21

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<210> 22

<211> 1019

<212> PRT

<213> Homo sapiens

<400> 22

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      20              25              30
Gly Gln Ala Pro Leu Asp Pro Val Tyr Leu Pro Ala Ala Leu Glu Leu
      35              40              45
Leu Asp Ala Pro Glu His Phe Arg Val Gln Gln Val Gly His Tyr Pro

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85	90	95
Phe Ala Thr Gln Gln Val Val Pro Pro Arg Val Thr Glu Pro His Gln		
100	105	110
Arg Pro Val Pro Trp Asp Val Arg Ala Val Ser Val Glu Ala Ala Val		
115	120	125
Thr Pro Ala Glu Pro Tyr Ala Arg Val Leu Phe His Leu Lys Gly Gln		
130	135	140
Asp Trp Pro Pro Gly Ser Gly Ser Leu Pro Cys Ala Arg Leu His Ala		
145	150	155
Thr His Pro Ala Gly Thr Ala His Gln Ala Cys Arg Phe Gln Pro Ser		
165	170	175
Leu Gly Ala Cys Val Val Glu Leu Glu Leu Pro Ser His Trp Phe Ser		
180	185	190
Gln Ala Ser Thr Thr Arg Ala Glu Leu Ala Tyr Thr Leu Glu Pro Ala		
195	200	205
Ala Glu Gly Pro Gly Gly Cys Gly Ser Gly Glu Glu Asn Asp Pro Gly		
210	215	220
Glu Gln Ala Leu Pro Val Gly Gly Val Glu Leu Arg Pro Ala Asp Pro		
225	230	235
Pro Gln Tyr Gln Glu Val Pro Leu Asp Glu Ala Val Thr Leu Arg Val		
245	250	255
Pro Asp Met Pro Val Arg Pro Gly Gln Leu Phe Ser Ala Thr Leu Leu		
260	265	270
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290	295	300
Trp Thr Ala Lys Leu Asp Arg Phe Lys Gly Ser Arg His His Thr Thr		
305	310	315
Leu Ile Thr Cys His Arg Ala Gly Leu Thr Glu Pro Asp Ser Ser Pro		
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Leu Glu Leu Ser Glu Phe Leu Trp Val Asp Phe Val Val Glu Asn Ser		

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Tyr Pro Gly Gln Ala Pro Glu Ala Glu Lys Asp Lys Met Val Trp Glu		
370	375	380
Ile Leu Val Ser Glu Arg Asp Ile Arg Ala Leu Ile Pro Leu Ala Lys		
385	390	395
Ala Glu Glu Leu Val Asn Thr Ala Pro Leu Thr Gly Val Pro Gln His		
405	410	415
Val Pro Val Arg Leu Val Thr Val Asp Gly Gly Gly Ala Leu Val Glu		
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Val Ser Glu Ala Cys Asp Ala Val Phe Val Ala Gly Lys Glu Ser Arg		
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Gly Ala Arg Gly Val Arg Val Asp Phe Trp Trp Arg Arg Leu Arg Ala		
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Ser Leu Arg Leu Thr Val Trp Ala Pro Leu Leu Pro Leu Arg Ile Glu		
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Leu Thr Asp Thr Thr Leu Glu Gln Val Arg Gly Trp Arg Val Pro Gly		
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Pro Ala Glu Gly Pro Ala Glu Pro Ala Ala Glu Ala Ser Asp Glu Ala		
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Glu Arg Arg Ala Arg Gly Cys His Leu Gln Tyr Gln Arg Ala Gly Val		
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Arg Phe Leu Ala Pro Phe Ala Ala His Pro Leu Asp Gly Gly Arg Arg		
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Leu Thr His Leu Leu Gly Pro Asp Trp Leu Leu Asp Val Ser His Leu		
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Val Ala Pro His Ala Arg Val Leu Asp Ser Arg Val Ala Ser Leu Glu		
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Gly Gly Arg Val Val Val Gly Arg Glu Pro Gly Val Thr Ser Ile Glu		
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Val Arg Ser Pro Leu Ser Asp Ser Ile Leu Gly Glu Gln Ala Leu Ala		
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Val Thr Asp Asp Lys Val Ser Val Leu Glu Leu Arg Val Gln Pro Val		

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Val Thr Ala Thr Cys Trp Ala Gln Ser Ala Leu Pro Ala Pro Lys Gln			
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Thr Gly Val Arg Gly Lys Phe Glu Arg Ala Glu Glu Glu Ala Arg Lys			
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Val Pro Ala Pro Gln His Val Thr Glu Leu Glu Leu Gly Met Tyr Ala			
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Leu Leu Gly Val Phe Cys Val Ala Ile Phe Ile Phe Leu Val Asn Gly			
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Val Val Phe Val Leu Arg Tyr Gln Arg Lys Glu Pro Pro Asp Ser Ala			
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Thr Asp Pro Thr Ser Pro Gln Pro His Asn Trp Val Trp Leu Gly Thr			
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<211> 3057

<212> DNA

<213> Homo sapiens

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<211> 3502

<212> DNA

<213> Homo sapiens

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<211> 1023

<212> PRT

<213> Homo sapiens

<400> 25

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 Gly His Tyr Pro Pro Ala Asn Ser Ser Leu Ser Ser Arg Ser Glu Thr
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 Phe Leu Leu Leu Gln Pro Trp Pro Arg Ala Gln Pro Leu Leu Arg Ala
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<212> DNA

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<212> PRT

<213> Homo sapiens

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